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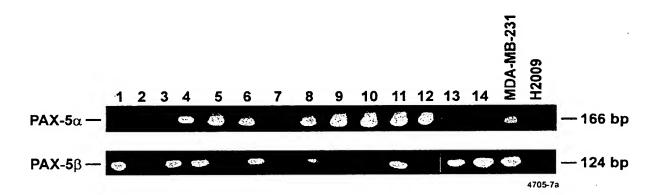
FIG. 1

Frequency of PAX5  $\alpha$  and  $\beta$  methylation in tumor cell line primary tumors and non-malignant specimens

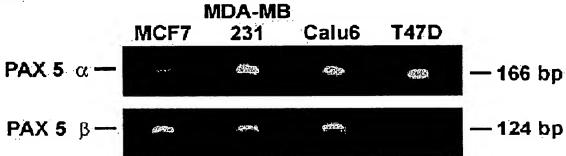
The second secon				
		Nun	Number Methylated (%)	(%)
Sample Type	Samples Assayed	Alpha	Beta	Alpha + Beta
Breast				
Tumor Cell Lines	10	6 (90)	(09) 9	(09) 9
Tumors	30	21 (70)	15 (50)	11 (37)
Lung				
Tumor Cell Lines	11	9 (82)	6 (55)	6 (55)
Tumors				
ADC	25	16 (64)	13 (52)	10 (40) <sup>c</sup>
SCC	23	17 (74)	14 (61)	$12(52)^c$
Colon				
Tumors	42	$6(21)^{d}$	14 (33) <sup>d</sup>	4 (9.5)
BECs"	35	1 (2.9)	1 (2.9)	0 (0)
Sputum"	39	5 (13)	1 (2.6)	0 (0)
Blood lymphocytes <sup>b</sup>	20	(0) 0	(0) (	. (0) 0
<sup>a</sup> Cancer-free heavy smokers		(2)	7212	

Cancer-free heavy smokers  $^{\prime}$  Cancer-free nonsmokers  $^{\prime}$  P=0.05 for association of PAX5  $\alpha$  methylation to  $\beta$  methylation in lung tumors.  $^{\prime}$  p<0.0001 as compared to methylation in lung or breast tumors.

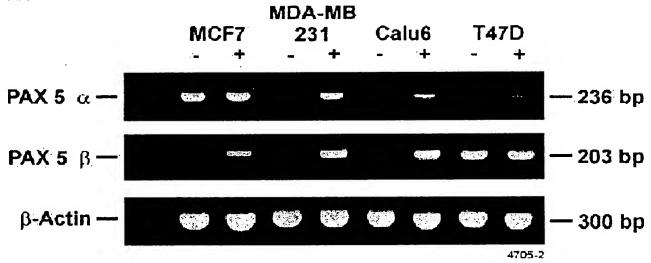
FIG. 2







## FIG. 3B.



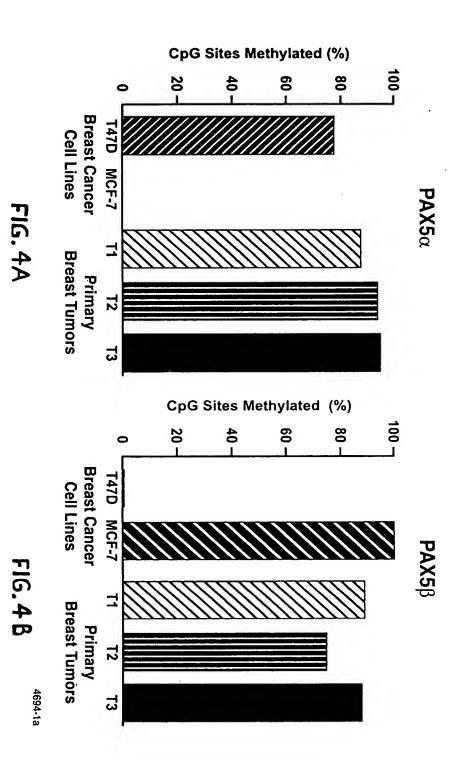


FIG. A

Frequency of methylation of the Novel Helix Loop Helix Binding Protein, Novel Gene 2, and Beta3 genes in tumor cell lines, primary and non-malignant specimens tumors

		Number Methylated (%)	
Sample Type	HLHBP	Novel Gene 2	Beta 3
Breast			
Tumor Cell Lines	8/10 (80)	5/8 (63)	(9) (9)
Tumors	NΩ	ND	6/15 (40)
Lung			
Tumor Cell Lines	8/11 (73)	1/7 (14)	1/12 (8)
Tumors			
ADC	ND	ND	11/21 (52)
SCC	13/22 (59)	ND	1/14 (7)
Blood lymphocytes	0/11 (0)	0/11(0)	0/11(0)

HLHBP, helix loop helix binding protein, ADC, adenocarcinoma; SCC, squamous cell carcinoma; BECs, bronchial epithelial cells "Cancer-free nonsmokers ND, not determined